Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L1	5057982	free energy, computational, structure-based, drug design, ligand, protein, binding, program, optimize	US-PGPUB; USPAT; DERWENT	OR	ON	2005/11/03 16:38
L2	228	free energy, computational, structure-based, drug design, ligand, protein, binding, program, optimize	US-PGPUB; USPAT; DERWENT	AND	ON	2005/11/03 16:38
L3	220	L2 and conformations	US-PGPUB; USPAT; DERWENT	AND	ON	2005/11/03 16:38
L4	217	L3 and structural information	US-PGPUB; USPAT; DERWENT	AND	ON	2005/11/03 16:38
L5	217	L4 and binding conformations	US-PGPUB; USPAT; DERWENT	AND	ON	2005/11/03 16:38
L6	214	L5 and binding region	US-PGPUB; USPAT; DERWENT	AND	ON	2005/11/03 16:38
L7	18	L5 and processor and solvation and model	US-PGPUB; USPAT; DERWENT	AND	ON	2005/11/03 16:39